



INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(51) International Patent Classification ⁶ : C12N 15/53, 15/29, 5/04, A01H 5/00, 1/00		A1	(11) International Publication Number: WO 96/37617
			(43) International Publication Date: 28 November 1996 (28.11.96)
(21) International Application Number: PCT/AU96/00310 (22) International Filing Date: 22 May 1996 (22.05.96) (30) Priority Data: PN 3098 23 May 1995 (23.05.95) AU PN 5600 26 September 1995 (26.09.95) AU (71) Applicant (for all designated States except US): COMMON-WEALTH SCIENTIFIC AND INDUSTRIAL RESEARCH ORGANISATION [AU/AU]; Limestone Avenue, Campbell, ACT 2601 (AU). (72) Inventor; and (75) Inventor/Applicant (for US only): ROBINSON, Simon, Piers [AU/AU]; 28 Jervois Street, Hawthorn, S.A. 5062 (AU). (74) Agent: PHILLIPS ORMONDE & FITZPATRICK; 367 Collins Street, Melbourne, VIC 3000 (AU).			(81) Designated States: AL, AM, AT, AU, AZ, BB, BG, BR, BY, CA, CH, CN, CZ, DE, DK, EE, ES, FI, GB, GE, HU, IS, JP, KE, KG, KP, KR, KZ, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, TJ, TM, TR, TT, UA, UG, US, UZ, VN, ARIPO patent (KE, LS, MW, SD, SZ, UG), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG). Published <i>With international search report.</i>
(54) Title: POLYPHENOL OXIDASE GENES FROM LETTUCE AND BANANA			
(57) Abstract The present invention provides methods for preparing nucleic acids encoding polyphenol oxidase (PPO), fragments and derivatives thereof. The present invention also provides nucleic acids encoding banana or lettuce PPO, or antisense to banana or lettuce PPO, fragments and derivatives thereof. Vectors including such nucleic acids, methods of using such nucleic acids and transgenic plants are also provided.			

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POLYPHENOL OXIDASE GENES FROM LETTUCE AND BANANA

The present invention relates to the isolation of genes encoding polyphenol oxidase (PPO) from plants.

5 Browning of plant tissues often occurs following injury or damage and this generally results in spoilage of fruit and vegetables. Undesirable browning also occurs during processing of plant materials to produce food or other products. Steps are taken during transport, storage, and processing to prevent these browning reactions. Often this involves the use of chemicals such as sulphur dioxide but the use of these substances is likely to be restricted in the future due to concerns about their safety and consumer acceptance. For example, the US Food and Drug Administration banned the use of sulphite for most fresh fruit and vegetables in 1986. The production of fruit and vegetable varieties with an inherently low susceptibility to brown would remove the need for these chemical treatments.

15 It will be understood that browning in plants is predominantly catalysed by the enzyme PPO. PPO is localised in the plastids of plant cells whereas the phenolic substrates of the enzyme are stored in the plant cell vacuole. This compartmentation prevents the browning reaction from occurring unless the plant cells are damaged and the enzyme and its substrates are mixed.

20 The prior art includes International Application PCT/AU92/00356 to the present applicant which describes the cloning of PPO genes from grapevine, broad bean leaf, apple fruit and potato tuber. This application recognises that PPO levels in plants may be manipulated by increasing or decreasing expression of PPO gene. The application also identifies two conserved copper binding sites in PPO genes, designated CuA and CuB. However, the method described in PCT/AU92/00356 which was used to clone the PPO genes from apple and potato involved the use of an oligo dT reverse primer for polymerase chain reaction (PCR). Whilst the method is acceptable, in some tissues, it does not give rise to a strong band of the predicted size or else it gives rise to many additional products making it difficult to resolve the PPO fragment.

30 Accordingly, it is an object of the present invention to overcome or at least alleviate one or more of the difficulties related to the prior art.

In a first aspect of the present invention there is provided a method for preparing nucleic acid encoding PPO, fragments and derivatives thereof, which method includes

providing

- 5 a source of a polypeptide having PPO activity,
 a first primer having a sequence corresponding to a first conserved
 region of PPO in sense orientation, and
 a second primer having a sequence corresponding to a second
 conserved region of PPO in antisense orientation;
10 isolating RNA from the source of polypeptide having PPO activity;
 treating the RNA to construct copy DNA (cDNA) therefrom; and
 amplifying the cDNA so formed using the first and second primers.

Applicant has found that the method of the present invention, which involves the use of a second primer based on PPO, means that there is less
15 likelihood that other (non-PPO) genes are amplified. Furthermore, the method of the present invention dramatically increases the amount of genuine product formed in most cases. Moreover, the added specificity provided by the second PPO-based primer makes it possible to clone PPO more readily from certain plants in which it was difficult to obtain a clone using one primer and oligo-dT.
20 For example, with lettuce cDNA the applicant saw only a faint smear of a range of products with GEN3/GEN8 and oligo-dT but strong bands of the predicted size with GEN3/GEN8 and REV1.

The terms "nucleic acid encoding banana/lettuce PPO" and "banana/lettuce PPO gene" as used herein should be understood to refer to a
25 banana/lettuce PPO gene or a sequence substantially homologous therewith. For example, these terms include sequences which differ from the specific sequences given in the Examples hereto but which, because of the degeneracy of the genetic code, encode the same protein. Applicants have found that there are families of PPO genes in most plants. Thus, there are likely to be other PPO
30 genes in lettuce and banana, in addition to those which have been isolated. These could be cloned using the methods of the present invention. Thus, the terms "nucleic acid encoding banana/lettuce PPO" and "banana/lettuce PPO

gene" should be understood to include banana/lettuce PPO genes other than those specific genes that have been isolated. The terms may also include presequences such as chloroplast transit sequence as well as sequences encoding mature PPO protein.

- 5 The term "derivative" as used herein includes nucleic acids that have been chemically or otherwise modified, for example mutated, or labelled, or nucleic acids incorporating a catalytic cleavage site.

 The term "fragment" includes functionally active fragments of a PPO gene which encode a polypeptide or peptide having PPO activity.

- 10 The source of polypeptide having PPO activity is preferably a source of polypeptide having banana or lettuce PPO activity. The source of polypeptide having banana PPO activity may be banana fruit, preferably young banana fruit, more preferably the flesh of young banana fruit. The source of polypeptide having lettuce PPO activity may be lettuce leaves, preferably young lettuce
15 leaves.

- The RNA may be isolated by any suitable method including extraction for example with a detergent such as CTAB, use of an oligo-dT spun column as described in PCT/AU92/00356 the entire disclosure of which is incorporated herein by reference, or use of a commercially available kit such as the
20 PolyATtract 1000 system from Promega Corporation.

 The step of treating the RNA to construct cDNA according to this aspect of the present invention may include

 treating the RNA with reverse transcriptase and an adapter primer to form cDNA.

- 25 The adapter primer may be an oligonucleotide adapter primer including the following sequence or part thereof:

5'-GACTCGAGTCGACATCGATTTTTTTTTTTTTTTT-3'

- The first primer has a sequence corresponding to a first conserved region of PPO. Preferably the first primer has a sequence corresponding to at least a
30 portion of or in close proximity to a first copper binding site of PPO. The second primer has a sequence corresponding to a second conserved region of PPO. Preferably the second primer has a sequence corresponding to at least a portion

of or in close proximity to a second copper binding site of PPO. More preferably the first primer has a sequence corresponding to at least a portion of or in close proximity to one of the CuA or CuB binding sites of PPO, and the second primer has a sequence corresponding to at least a portion of or in close proximity to the other of the CuA or CuB binding sites of PPO.

The first and second primers may be degenerate. The first primer may include one of the following sequences or part thereof:

5'-GCGAATTCTT[TC][TC]TICCITT[TC]CA[TC][AC]G-3'

5'-GCGAATTCGATCCACATT[TC]GC[GT]TTICC-3'.

The second primer may include the following sequence or part thereof

5'-GCCTGCAGCCACATC[AG][AG]TCIAC[AG]TT-3'.

The cDNA may be amplified using the polymerase chain reaction (PCR).

Those skilled in the art will appreciate that if the Cu binding sites are internal, the nucleic acid isolated will be a fragment of the PPO gene lacking 3' and 5' termini. However, it is possible to determine the complete nucleic acid sequence of the PPO gene and to prepare or isolate nucleic acid encoding such PPO or antisense to such PPO.

Accordingly, in a further aspect of the present invention there is provided a method for preparing nucleic acid encoding the 3' end of PPO, which method includes

providing

a source of polypeptide having PPO activity

a primer in sense orientation; and

an adapter primer;

isolating RNA from the source of polypeptide having PPO activity;

treating the RNA to construct cDNA therefrom; and

amplifying the cDNA so formed using the primers.

In a further aspect of the present invention there is provided a method for preparing nucleic acid encoding the 5' end of PPO, which method includes

providing

a source of polypeptide having PPO activity,

an anchor,

- primers in antisense orientation; and
an anchor primer;
isolating RNA from the source of polypeptide having PPO activity;
treating the RNA to construct cDNA therefrom;
5 attaching the anchor to the 5' end of the cDNA so formed; and
amplifying the cDNA using the primers.

The source of polypeptide having PPO activity is preferably a source of polypeptide having banana or lettuce PPO activity. The source of polypeptide having banana PPO activity may be banana fruit, preferably young banana fruit,
10 more preferably the flesh of young banana fruit. The source of polypeptide having lettuce PPO activity may be lettuce leaves, preferably young lettuce leaves.

The RNA may be isolated by any suitable method including extraction for example with a detergent such as CTAB, use of an oligo-dT spun column as
15 described in PCT/AU92/00356 the entire disclosure of which is incorporated herein by reference, or use of a commercially available kit such as the PolyATtract 1000 system from Promega Corporation.

The step of treating the RNA to construct cDNA according to this aspect of the present invention may include
20 treating the RNA with reverse transcriptase and an adapter primer to form cDNA.

The adapter primer may be an oligonucleotide adapter primer including the following sequence or part thereof:

5'-GACTCGAGTCGACATCGATTTTTTTTTTTTTTTT-3'

25 The primer in sense orientation may be a lettuce PPO specific primer. The primer in sense orientation may include the following sequence or part thereof:

5'-CGCTGGGTGGGTAATTCTAGGATG-3'.

The primer in sense orientation may be a banana PPO specific primer. The primer in sense orientation may include the following sequence or part
30 thereof:

5'-AGTCATCCACAATGCGGCGCACATG-3'.

The adapter primer may include the following sequence or part thereof:

5'-GACTCGAGTCGACATCG-3'.

The primers in antisense orientation may be lettuce PPO specific primers.

5 The primers in antisense orientation may include the following sequences or part thereof:

5'-TGCTGTTCTGTTCGAACATGGCAG-3'

5'-TATACAAGTGGCACCAGTGTCTGC-3'.

The primers in antisense orientation may be banana PPO specific primers.

10 The primers in antisense orientation may include the following sequences or part thereof:

5'-CCGCATTGTGGATGACTTCCATCTG-3'

5'-CCAGAATGGGATGGTGAAGGTGTCTG-3'.

15 The anchor may be of any suitable type. The anchor may be attached by ligation for example using T4 RNA ligase. The anchor primer should be capable of hybridizing with the anchor.

The cDNA may be amplified using PCR.

Those skilled in the art will appreciate that using the methods of the present invention it is possible to determine the complete nucleic acid sequence of the PPO gene of interest and to prepare or isolate nucleic acid encoding such
20 PPO or antisense to such PPO.

In a further aspect of the present invention, there is provided a nucleic acid encoding banana PPO or antisense to banana PPO, fragments and derivatives thereof. Preferably the nucleic acid has the sequence shown in Fig. 2 or Fig. 3, fragments and derivatives thereof, and substantially homologous sequences.

25 In a further aspect of the present invention, there is provided a nucleic acid encoding lettuce PPO or antisense to lettuce PPO, fragments and derivatives thereof. Preferably the nucleic acid has the sequence shown in Fig. 1, fragments and derivatives thereof, and substantially homologous sequences.

The nucleic acid may be prepared by a method as hereinbefore described.

30 The nucleic acid may be modified, for example by inclusion of a catalytic cleavage site.

In a further aspect of the present invention there is provided a method for preparing a recombinant vector including a nucleic acid encoding banana PPO or antisense to banana PPO, fragments and derivatives thereof, which method includes

5 providing

nucleic acid encoding banana PPO or antisense to banana PPO,
fragments and derivatives thereof; and
a vector; and

10 reacting the nucleic acid and the vector to deploy the nucleic acid within
the vector.

In a further aspect of the present invention there is provided a method for preparing a recombinant vector including a nucleic acid encoding lettuce PPO or antisense to lettuce PPO, fragments and derivatives thereof, which method includes

15 providing

nucleic acid encoding lettuce PPO or antisense to lettuce PPO,
fragments and derivatives thereof; and
a vector; and

20 reacting the nucleic acid and the vector to deploy the nucleic acid within
the vector.

The nucleic acid may be prepared by a method as hereinbefore described.

The nucleic acid may be modified, for example by inclusion of a catalytic cleavage site.

25 The vector may be a plasmid expression vector. For example Bluescript SK⁺ has been found to be suitable. Alternatively, the vector may be a binary vector. The recombinant vector may contain a promoter, preferably a constitutive promoter upstream of the nucleic acid.

The cloning step may take any suitable form. A preferred form may include

30 fractionating the cDNA, for example on a column or a gel;

isolating a fragment of the expected size, for example from the column or gel; and

ligating said fragment into a suitable restriction enzyme site of the vector, for example the EcoRV site of a Bluescript SK⁺ vector.

In order to test the clones so formed, a suitable microorganism may be transformed with the vector, the microorganism cultured and the polypeptide encoded therein expressed. The microorganism may be a strain of Escherichia coli, for example E.coli DH5 has been found to be suitable. Alternatively, appropriate vectors may be used to transform plants.

In a further aspect of the present invention there is provided a recombinant vector including a nucleic acid encoding banana PPO or antisense to banana PPO, fragments and derivatives thereof, which vector is capable of being replicated, transcribed and translated in a unicellular organism or alternatively in a plant.

In a further aspect of the present invention there is provided a recombinant vector including a nucleic acid encoding lettuce PPO or antisense to lettuce PPO, fragments and derivatives thereof, which vector is capable of being replicated, transcribed and translated in a unicellular organism or alternatively in a plant.

The nucleic acid may be prepared by a method as hereinbefore described.

The nucleic acid may be modified, for example by inclusion of a catalytic cleavage site.

The vector may be a plasmid expression vector. For example Bluescript SK⁺ has been found to be suitable. Alternatively, the vector may be a binary vector. The recombinant vector may contain a promoter, preferably a constitutive promoter upstream of the nucleic acid encoding banana PPO or antisense to banana PPO, fragments and derivatives thereof.

The microorganism may be a strain of Escherichia coli, for example E.coli DH5 has been found to be suitable.

In a further aspect of the present invention there is provided a method of decreasing the level of PPO activity in a plant tissue, which method includes providing

a nucleic acid encoding banana PPO, a modified nucleic acid encoding banana PPO, or a nucleic acid antisense to banana PPO, fragments and derivatives thereof; and

a plant sample; and
introducing said nucleic acid into said plant sample to produce a transgenic plant.

In a further aspect of the present invention there is provided a method of
5 decreasing the level of PPO activity in a plant tissue, which method includes
providing

a nucleic acid encoding lettuce PPO, a modified nucleic acid
encoding lettuce PPO, or a nucleic acid antisense to lettuce PPO,
fragments and derivatives thereof; and

10 a plant sample; and

introducing said nucleic acid into said plant sample to produce a transgenic plant.

PPO activity may be decreased by the use of sense constructs (cosuppression). Alternatively the nucleic acid may include a sequence encoding
15 antisense mRNA to banana or lettuce PPO or a functionally active fragment thereof. Alternatively the nucleic acid may encode banana or lettuce PPO or a functionally active fragment thereof and incorporate a catalytic cleavage site (ribozyme). The nucleic acid may be included in a recombinant vector as hereinbefore described. In a preferred aspect, the nucleic acid may be included
20 in a binary vector. In a further preferred aspect, the introduction of a binary vector into the plant may be by infection of the plant with an Agrobacterium containing the binary vector or by bombardment with nucleic acid coated microprojectiles. Methods for transforming banana with Agrobacterium are known to those skilled in the art and are described in, for example, May et al.,
25 Bio/technology (1995) 13:486-492, the entire disclosure of which is incorporated herein by reference. Methods for transforming banana by bombardment with DNA coated microprojectiles are known to those skilled in the art and are described in, for example, Sagi et al., Bio/technology (1995) 13:481-485, the entire disclosure of which is incorporated herein by reference. Methods for
30 transforming lettuce using Agrobacterium are known to those skilled in the art and are described in, for example, Michelmores et al., Plant Cell Reports (1987) 6:439-442, and Curtis et al., Journal of Experimental Botany (1994)

45:1141-1149.

In a further aspect of the present invention there is provided a method of increasing the level of PPO activity in a plant tissue, which method includes providing

- 5 a nucleic acid encoding banana PPO or a fragment thereof; and
 a plant sample; and

introducing said nucleic acid into said plant sample to produce a transgenic plant.

- In a further aspect of the present invention there is provided a method of
10 increasing the level of PPO activity in a plant tissue, which method includes
 providing

 a nucleic acid encoding lettuce PPO or a fragment thereof; and
 a plant sample; and

- introducing said nucleic acid into said plant sample to produce a
15 transgenic plant.

The nucleic acid may be included in a recombinant vector as hereinbefore described. In a preferred aspect, the nucleic acid may be included in a binary vector. In a further preferred aspect, the introduction of the binary vector into the plant may be by infection of the plant with an Agrobacterium containing the binary
20 vector or by bombardment with nucleic acid coated microprojectiles.

The plant may be of any suitable type. However the method is particularly applicable to banana or lettuce.

- In a further aspect of the present invention there is provided a transgenic plant, which plant contains nucleic acid capable of modifying expression of the
25 normal banana PPO gene.

The plant may be of any suitable type. Preferably, the plant is banana.

In a further aspect of the present invention there is provided a transgenic plant, which plant contains nucleic acid capable of modifying expression of the normal lettuce PPO gene.

- 30 The plant may be of any suitable type. Preferably, the plant is lettuce.

The nucleic acid may be as hereinbefore described.

In a still further aspect of the present invention there is provided a plant vaccine including nucleic acid encoding banana PPO or antisense to banana PPO, fragments and derivatives thereof.

5 In a still further aspect of the present invention there is provided a plant vaccine including nucleic acid encoding lettuce PPO or antisense to lettuce PPO, fragments and derivatives thereof.

The present invention will now be more fully described with reference to the accompanying Examples. It should be understood, however, that the description following is illustrative only and should not be taken in any way as a
10 restriction on the generality of the invention described above.

In the Figures:

FIGURE 1: The composite LPO1 cDNA nucleotide sequence and derived protein sequence encoding both the putative chloroplast transit sequence and the mature lettuce PPO protein.

15 FIGURE 2: The BANPPO1 cDNA nucleotide sequence and derived protein sequence encoding both the putative chloroplast transit sequence and the mature banana PPO protein.

FIGURE 3: The BANPPO11 cDNA nucleotide sequence and derived protein sequence encoding part of a banana PPO protein.

20

EXAMPLE 1

Cloning Lettuce PPO Genes

Messenger RNA (mRNA) was isolated directly from young leaves of lettuce using the PolyATtract 1000 system from Promega Corporation. First strand cDNA was synthesised with reverse transcriptase using a Timesaver
25 cDNA Synthesis Kit (Pharmacia Biotech) utilising an oligo-dT primer adapter as described in Frohman, MA (1990) in "PCR Protocols : A Guide to Methods and Applications" (MA Innis, DH Gelfrand, JJ Sninsky and TJ White, eds) Academic Press, New York pp 28-38, the entire disclosure of which is incorporated herein by reference:

30

B26 : (5'-GACTCGAGTCGACATCGATTTTTTTTTTTTTTTT-3').

Oligonucleotide primers were designed based on known plant PPO DNA sequences in the conserved regions of the gene which encode the copper binding sites, CuA and CuB as described in Dry, IB and Robinson, SP (1994) "Molecular cloning and characterisation of grape berry polyphenol oxidase", Plant
5 Molecular Biology 26 : 495-502, the entire disclosure of which is incorporated herein by reference. Two forward primers designed around the CuA site (GEN3 and GEN8) and one reverse primer designed around the CuB site (REV1) were synthesised:

GEN3 : (5'-GCGAATTCTT[TC][TC]TICCITT[TC]CA[TC][AC]G-3')
10 GEN8 : (5'-GCGAATTCGATCCACATT[TC]GC[GT]TTICC-3')
REV1 : (5'-GCCTGCAGCCACATC[AG][AG]TCIAC[AG]TT-3')

Although the primers are in the region of the Cu binding sites, one of them (GEN8) is just outside of what is traditionally accepted to be a Cu binding site of the enzyme.

15 The first strand cDNA was amplified by the polymerase chain reaction (PCR) essentially according to the method of Frohman using GEN3 and REV1 or GEN8 and REV1 primers, each at a final concentration of 1 μ M (Dry et al.). Amplification involved an initial program of 2 cycles of denaturation at 94°C for 1 min, annealing at 37°C for 2 min, a slow ramp to 72°C over 2 min and elongation
20 at 72°C for 3 min, followed by 25 cycles of denaturation at 94°C for 1 min, annealing at 55°C for 1 min, and elongation at 72°C for 3 min. A sample of the amplified DNA was run on an agarose gel and stained with ethidium bromide to determine the size of the PCR products and the remainder was purified and concentrated using PCR Wizard Prep columns (Promega Corporation).

25 The purified DNA was cloned into Eco RV-cut Bluescript SK⁺ vector (Stratagene) which had been T-tailed with Taq Polymerase and the ligated DNA was introduced into E.coli DH5 α by electroporation. Recombinant clones which had an insert of the predicted size were selected and their DNA sequence was determined by automated sequencing. Three putative lettuce PPO clones
30 (LPO316, LPO812 and LPO813) were identified based on their homology to known plant PPO genes.

Using this sequence information a specific forward primer (LET3P) and two reverse primers (LET5P1 and LET5P2) were synthesised:

LET3P : (5'-CGCTGGGTGGGTAATTCTAGGATG'3-)

LET5P1 : (5'-TGCTGTTCTGTTCTGAACATGGCAG-3') ,

5 LET5P2 : (5'-TATACAAGTGGCACCAGTGTCTGC-3')

To obtain the 3'-end of the lettuce PPO gene, the first strand cDNA described above was amplified by the same PCR procedure using 1 μ M LET3P primer and 100 nM adapter primer:

B25 : (5'-GACTCGAGTCGACATCG-3').

10 The amplified cDNA was purified as described above and run on a 2% Nusieve GTG (FMC Bioproducts) agarose gel. A 1000bp fragment was excised from the gel and the DNA was cloned into T-tailed, Eco RV-cut Bluescript SK⁺ to yield the 3'- end clones LPO9 and LPO10, which were sequenced.

The 5'-end of the lettuce PPO gene was cloned by a modification of the 5'-
15 RACE procedure originally described by Frohman using a 5'-AmpliFINDER RACE kit (Clontech Laboratories). First strand cDNA was synthesised from mRNA with reverse transcriptase using the LET5P2 primer and an AmpliFINDER anchor was ligated onto the 5'-end of the cDNA. The cDNA was amplified by PCR with LET5P1 primer and the AmpliFINDER anchor primer. The amplified cDNA was
20 purified as described above and run on a 2% Nusieve GTG (FMC Bioproducts) agarose gel. An 850bp fragment was excised from the gel and the DNA was cloned into T-tailed Eco RV-cut Bluescript SK⁺ to give the 5'-end clones LPO4, LPO5, LPO6, and LPO7, which were sequenced.

The 5'- and 3'-clones were found to have the predicted overlapping
25 sequences with the original clone and the complete sequence of lettuce PPO (LPO1) was derived by combining the sequences from the various clones (Figure 1).

EXAMPLE 2**Cloning Banana PPO Genes**

Total RNA was isolated from young banana fruit. Fruit tissue (3g) was frozen and ground to a fine powder in liquid nitrogen with a coffee grinder then
5 added to 20 ml of extraction buffer (2% hexadecyltrimethylammonium bromide (CTAB), 2% polyvinyl pyrrolidone, 100 mM Tris-HCl, pH 8.0, 25 mM EDTA, 2 M NaCl, 0.05% spermidine, 2% β -mercaptoethanol) at 65°C. The extract was mixed with 20 ml of chloroform / IAA then centrifuged for 20 minutes at 5,000 RPM and the aqueous phase was re-extracted with chloroform / IAA. The aqueous phase
10 was filtered through Miracloth and 0.25 volumes of 10 M LiCl were added then the sample was incubated overnight at 4°C before centrifuging for 20 minutes at 8,000 RPM. The supernatant was removed and the pellet was resuspended in 0.5 ml of 1 M NaCl, 0.5% SDS, 10 mM Tris, pH 8.0, 1 mM EDTA. The RNA was extracted once with an equal volume of chloroform / IAA and 2 volumes of
15 ethanol was added. After incubation for 40 mins at -70°C the solution was centrifuged for 15 minutes at 10,000 RPM. The supernatant was removed and the pellet was rinsed with 80% ethanol, drained, and dried. The pellet was resuspended in 50 μ l of sterile water.

First strand cDNA was synthesised from 10 μ g total RNA with reverse
20 transcriptase as described in Dry, I.B. and Robinson, S.P. (1994) "Molecular cloning and characterisation of grape berry polyphenol oxidase", Plant Molecular Biology 26 : 495-502, the entire disclosure of which is incorporated herein by reference, utilising an oligo-dT primer adapter (Frohman, M.A. (1990) in "PCR
25 Protocols : A Guide to Methods and Applications" (M.A. Innis, D.H. Gelfrand, J.J. Sninsky and T.J. White, eds.) Academic Press, New York pp 28-38, the entire disclosure of which is incorporated herein by reference) :

B26 : (5'-GACTCGAGTCGACATCGATTTTTTTTTTTTTTTTTTTT-3')

Oligonucleotide primers were designed based on known plant PPO DNA sequences in the conserved regions of the gene which encode the copper binding
30 sites, CuA and CuB (Dry et al.). A forward primer designed around the CuA site (GEN3) and a reverse primer designed around the CuB site (REV1) were synthesised :

GEN3 : (5'-GCGAATTCTT[TC][TC]TICCITT[TC]CA[TC][AC]G-3')

REV1 : (5'-GCCTGCAGCCACATIC[AG][AG]TCIAC[AG]TT-3')

The first strand reaction was amplified by the polymerase chain reaction (PCR) essentially according to the method of Frohman using GEN3 and REV1 primers, each at a final concentration of 1 μ M (Dry et al.). Amplification involved an initial program of 2 cycles of denaturation at 94° C for 1 min, annealing at 37° C for 2 min, a slow ramp to 72° C over 2 min and elongation at 72° C for 3 min, followed by 25 cycles of denaturation at 94° C for 1 min, annealing at 55° C for 1 min, and elongation at 72° C for 3 min. A sample of the amplified DNA was run on an agarose gel and stained with ethidium bromide to determine the size of the PCR products and the remainder was purified and concentrated using PCR Wizard Prep columns (Promega Corporation).

The purified DNA was cloned into Eco RV-cut Bluescript SK⁺ vector (Stratagene) which had been T-tailed with Taq Polymerase and the ligated DNA was introduced into *E. coli* DH5 α by electroporation. Recombinant clones which had an insert of the predicted size were selected and their DNA sequence was determined by automated sequencing. A putative banana PPO clone (BPO3) was identified based on its homology to known plant PPO genes.

Using this sequence information a specific forward primer (BAN1) and two specific reverse primers (BAN2R and BAN3R) were synthesised:

BAN 1 : (5'-AGTCATCCACAATGCGGCGCACATG-3')

BAN2R : (5'-CCGCATTGTGGATGACTTCCATCTG-3')

BAN3R : (5'-CCAGAATGGGATGGTGAAGGTGTGCG-3')

To obtain the 3'-end of this banana PPO gene, the first strand cDNA described above was amplified by the same PCR procedure using 1 μ M BAN1 primer and 100nM adapter primer:

B25 : (5'-GACTCGAGTCGACATCG-3')

The DNA was amplified using 25 cycles of denaturation at 94°C for 1 min, annealing at 55°C for 1 min, and elongation at 72°C for 3 min. The amplified DNA was purified using a QIAquick Spin PCR Purification Kit (QIAGEN) and run on a 2% Nusieve GTG (FMC Bioproducts) agarose gel. A 1000bp fragment was excised from the gel and the DNA was cloned into T-tailed Eco RV-cut Bluescript

SK⁺ to yield the 3'-end clone BPO17, which was sequenced and shown to encode the 3'-end of BPO3.

The 5'-end of BPO3 was cloned by a modification of the 5'-RACE procedure originally described by Frohmann. First strand cDNA was synthesised from
5 banana fruit RNA as described above but utilising the banana PPO specific primer BAN2R. The DNA was tailed with Terminal transferase as described in Frohmann and amplified by PCR with BAN3R and B26 primers, each at a final concentration of 1 μ M. The DNA was amplified using 30 cycles of denaturation at 94°C for 1 min, annealing at 55°C for 1 min, and elongation at 72°C for 3 min.
10 The amplified DNA was run on a 1.8% Nusieve GTG (FMC Bioproducts) agarose gel and a 700bp fragment was excised from the gel. The DNA was extracted with a QIAquick Gel Extraction Kit and cloned into T-tailed Eco RV-cut Bluescript SK⁺ to yield the 5'-end clone BPO26 which was sequenced and shown to encode the 5'-end of BPO3.

15 The overlapping clones BPO3, BPO17 and BPO26 were fully sequenced in both directions and the sequence of this banana PPO gene (BANPPO1) was derived by combining the sequences (Figure 2).

In the course of these experiments a number of clones were obtained from the banana fruit cDNA by PCR amplification using the B25 primer with one of the
20 degenerate primers based on conserved sequences in other plant PPO genes:

GEN7 : (5'-GCGAATTCAA[TC]GTIGA[TC][AC]GIATGTGG-3')

using the methods described above. Most of these clones were identical to BANPPO1 but one clone, designated BANPPO11, was found to be distinctly different and its sequence is shown in Figure 3.

25 Finally, it is to be understood that various other modifications and/or alterations may be made without departing from the spirit of the present invention as outlined herein.

CLAIMS:

1. A method for preparing nucleic acid encoding PPO, fragments and derivatives thereof, which method includes
providing
5 a source of a polypeptide having PPO activity,
a first primer having a sequence corresponding to a first conserved region of PPO in sense orientation, and
a second primer having a sequence corresponding to a second conserved region of PPO in antisense orientation;
10 isolating RNA from the source of polypeptide having PPO activity;
treating the RNA to construct copy DNA (cDNA) therefrom; and
amplifying the cDNA so formed using the first and second primers.
2. A method according to claim 1 wherein the first primer has a sequence
15 corresponding to at least a portion of or in close proximity to a first copper (Cu) binding site of PPO and the second primer has a sequence corresponding to at least a portion of or in close proximity to a second Cu binding site of PPO.
3. A method according to claim 2 wherein the nucleic acid encodes banana
20 or lettuce PPO and the source of polypeptide having PPO activity is a source of polypeptide having banana or lettuce PPO activity, respectively.
4. A method according to claim 3 wherein the first primer includes one of the following sequences or part thereof:
25 5'-GCGAATTCTT[TC][TC]TICCITT[TC]CA[TC][AC]G-3'
5'-GCGAATTCGATCCACITT[TC]GC[GT]TTICC-3'.
5. A method according to claim 4 wherein the second primer includes the following sequence or part thereof:
30 5'-GCCTGCAGCCACATIC[TG][AG]TCIAC[AG]TT-3'.

6. A method according to claim 3 wherein the step of treating the RNA to construct cDNA includes

treating the RNA with reverse transcriptase and an adapter primer including the following sequence or part thereof:

5 5'-GACTCGAGTCGACATCGATTTTTTTTTTTTTTTT-3'

to form cDNA.

7. A method according to claim 2, which method further includes providing

10 a source of polypeptide having PPO activity,
a primer in sense orientation, and
an adapter primer;

isolating RNA from the source of polypeptide having PPO activity;

treating the RNA to construct cDNA therefrom; and

15 amplifying the cDNA so formed using the primers to prepare nucleic acid
encoding the 3' end of PPO.

8. A method according to claim 7, which method further includes providing

20 a source of polypeptide having PPO activity,
an anchor,
primers in antisense orientation, and
an anchor primer;

isolating RNA from the source of polypeptide having PPO activity;

25 treating the RNA to construct cDNA therefrom;

attaching the anchor to the 5' end of the cDNA so formed; and

amplifying the cDNA using the primers to prepare nucleic acid encoding
the 5' end of PPO.

30 9. A method according to claim 8 wherein the nucleic acid encodes lettuce
PPO, the primer sense orientation includes the following sequence or part
thereof:

5'-CGCTGGGTGGGTAATTCTAGGATG-3', and

the primers in antisense orientation include the following sequences or part thereof:

5'-TGCTGTTCTGTTCTGAACATGGCAG-3'

5

5'-TATACAAGTGGCACCAGTGTCTGC-3'.

10. A method according to claim 8 wherein the nucleic acid encodes banana PPO, the primer in sense orientation includes the following sequence or part thereof:

10

5'-AGTCATCCACAATGCGGCGCACATG-3', and

the primers in antisense orientation include the following sequences or part thereof:

5'-CCGCATTGTGGATGACTTCCATCTG-3'

5'-CCAGAATGGGATGGTGAAGGTGTCG-3'.

15

11. A method according to claim 8 wherein the adapter primer includes the following sequence or part thereof:

5'-GACTCGAGTCGACATCG-3'.

20 12. A nucleic acid encoding banana PPO or antisense to banana PPO, fragments and derivatives thereof.

13. A nucleic acid according to claim 12 including a catalytic cleavage site.

25 14. A nucleic acid according to claim 12, having the sequence shown in Fig. 2 or Fig. 3, fragments and derivatives thereof, and substantially homologous sequences.

30 15. A nucleic acid encoding lettuce PPO or antisense to lettuce PPO, fragments and derivatives thereof.

16. A nucleic acid according to claim 15 including a catalytic cleavage site.

17. A nucleic acid according to claim 15, having the sequence shown in Fig. 1, fragments and derivatives thereof, and substantially homologous sequences.
18. A recombinant vector including a nucleic acid according to claim 12, which
5 vector is capable of being replicated, transcribed and translated in a unicellular organism or alternatively in a plant.
19. A recombinant vector including a nucleic acid according to claim 15, which
10 vector is capable of being replicated, transcribed and translated in a unicellular organism or alternatively in a plant.
20. A method of decreasing the level of PPO activity in a banana plant tissue, which method includes
providing
15 a nucleic acid according to claim 12; and
a plant sample; and
introducing said nucleic acid into said plant sample to produce a transgenic banana plant.
- 20 21. A method of decreasing the level of PPO activity in a lettuce plant tissue, which method includes
providing
a nucleic acid according to claim 15; and
a plant sample; and
25 introducing said nucleic acid into said plant sample to produce a transgenic lettuce plant.
22. A transgenic banana plant, which plant contains nucleic acid capable of modifying expression of the normal banana PPO gene.
30
23. A transgenic lettuce plant, which plant contains nucleic acid capable of modifying expression of the normal lettuce PPO gene.

1 GACCACCCATAGATGATGGCTTCTCTCGCCTTGTCTAGTCTTCCACCTCCACCACAACC 60
-----+-----+-----+-----+-----+
CTGGTGGGTATCTACTACCGAAGAGAGCGGAACAGATCAGAAGGGTGGAGGTGGTGTGG
M A S L A L S S L P T S T T T -

61 AAAAAACCCCTTATTTTCCAAAACATCCTCGCATGTTAAGCCATTCCATCGCTTCAAAGTT 120
-----+-----+-----+-----+-----+
TTTTTTGGGAATAAAAGGTTTTGTAGAGCGTACAATTCGGTAAGGTAGCGAAGTTTCAA
K K P L F S K T S S H V K P F H R F K V -

121 TCATGCAATGCACCCGCTGATAACAATGACAAAACCGTCAATAATTCTGATACCCCAAAG 180
-----+-----+-----+-----+-----+
AGTACGTTACGTGGGCGACTATTGTTACTGTTTTGGCAGTTATTAAGACTATGGGGTTTC
S C N A P A D N N D K T V N N S D T P K -

181 CTCATACTACCCAAAACACCCTTGAAACGCAGAACGTAGACAGGAGAACTTGCTTCTG 240
-----+-----+-----+-----+-----+
GAGTATGATGGGTTTTGTGGTGAACCTTGCCTCTTGCATCTGTCTCTTTGAACGAAGAC
L I L P K T P L E T Q N V D R R N L L L -

241 GGACTCGGAGGTCTCTACGGCGCTGCCAACTTGACGACCATTCCGTCAGCCTTTGGCATT 300
-----+-----+-----+-----+-----+
CCTGAGCCTCCAGAGATGCCGCGACGGTTGAACTGCTGGTAAGGCAGTCGGAACCGTAA
G L G G L Y G A A N L T T I P S A F G I -

301 CCCATCGCTGCTCCAGACAATATTTTCAGACTGTGTTGCTGCGACTTCAAACCTAAGGAAC 360
-----+-----+-----+-----+-----+
GGGTAGCGACGAGGTCTGTTATAAAGTCTGACACAACGACGCTGAAGTTTGGATTCTTTCG
P I A A P D N I S D C V A A T S N L R N -

361 AGCAAAGACGCTATAAGGGGACTAGCGTGTGTCTCCGGTGCTTTCAACAAACAAACCA 420
-----+-----+-----+-----+-----+
TCGTTTCTGCGATATTCCTGATCGCACAAACAGGAGGCCACGAAAGTTGTTTGTGGT
S K D A I R G L A C C P P V L S T N K P -

421 ATGGATTACGTCTTCTTCAAACCTGTGATTCTGTGTTTCGACCAGCTGCACAGAAAGCC 480
-----+-----+-----+-----+-----+
TACCTAATGCAGGAAGGAAGTTTGGGACACTAAGCACAAGCTGGTCGACGTGTCTTTCGG
M D Y V L P S N P V I R V R P A A Q K A -

481 ACTGCCGATTACACTGCTAAGTATCAACAAGCAATTCAAGCCATGAAGGATCTCCCCGAG 540
-----+-----+-----+-----+-----+
TGACGGCTAATGTGACGATTCATAGTTGTTTCGTTAAGTTCGGTACTTCCTAGAGGGGCTC
T A D Y T A K Y Q Q A I Q A M K D L P E -

541 GACCACCCACATAGCTGGAAGCAACAAGGAAGATTCACTGTGCTTATTGCAACGGTGGT 600
-----+-----+-----+-----+-----+
CTGGTGGGTGATCGACCTTCGTTGTTCCGTTCTAAGTGACACGAATAACGTTGCCACCA
D H P H S W K Q Q G K I H C A Y C N G G -

601 TACAATCAAGAACAAGTGGTTACCCGAATTTACAACCTTCAGATTACAACTCATGGCTC 660
-----+-----+-----+-----+-----+
ATGTTAGTTCTTGTTCACCAATGGGCTTAAATGTTGAAGTCTAAGTGTGAGTACCGAG
Y N Q E Q S G Y P N L Q L Q I H N S W L -

661 TTCTTTCTTTCCACCGGTGGTACCTCTATTCTACGAGAAGATATTGGGGAAGTTGATT 720
-----+-----+-----+-----+-----+
AAGAAAGGAAGGTGGCCACCATGGAGATAAAGATGCTCTTCTATAACCCCTTCAACTAA
F F P F H R W Y L Y F Y E K I L G K L I -

721 AATGATCCAACCTTTCGCTCTACCTTACTGGAACCTGGGATAACCCCTACTGGAATGGTTATT 780
-----+-----+-----+-----+-----+
TTACTAGGTTGAAAGCGAGATGGAATGACCTTGACCCTATTGGGATGACCTTACCAATAA
N D P T F A L P Y W N W D N P T G M V I -

781 CCTGCCATGTTTCGAACAGAACAGCAAACTAACTCTCTGTTTGACCCTTTAAGGGATGCG 840
-----+-----+-----+-----+-----+
GGACGGTACAAGCTTGTCTTGTGCTTTGATTGAGAGACAACTGGGAAATTCCTACGC
P A H F E Q N S K T N S L F D P L R D A -

FIG 1

841 AAACACCTCCACCTTCTATCTTTGATGTTGAATATGCTGGTGCAGACACTGGTGCCACT 900
-----+-----+-----+-----+-----+-----+
TTTGTGGAGGGTGGGAAGATAGAACTACAACCTTATACGACCACGTCTGTGACCACGGTGA
K H L P P S I F D V E Y A G A D T G A T -

901 TGTATAGACCAGATAGCCATTAATCTGTCTTCAATGTACAGACAGATGGTCACCAACTCC 960
-----+-----+-----+-----+-----+-----+
ACATATCTGGTCTATCGGTAATTAGACAGAAGTTACATGTCTGTCTACCAGTGGTTGAGG
C I D Q I A I N L S S M Y R Q M V T N S -

961 ACTGATACAAAACGATTCTTCGGTGGCGAATTTGTAGCTGGAAATGACCCTCTTGCGAGC 1020
-----+-----+-----+-----+-----+-----+
TGACTATGTTTTGCTAAGAAGCCACCGCTTAAACATCGACCTTTACTGGGAGAACGCTCG
T D T K R F F G G E F V A G N D P L A S -

1021 GAGTTCAACGTAGCTGGGACCGTAGAAGCTGGGGTTTCACACTGCGGCTCACCGCTGGGTG 1080
-----+-----+-----+-----+-----+-----+
CTCAAGTTGCATCGACCCTGGCATCTTCGACCCCAAGTGTGACGCCGAGTGGCGACCCAC
E F N V A G T V E A G V H T A A H R W V -

1081 GGTAATCTAGGATGGCCAACAGCGAAGACATGGGGAACCTTCTACTCCGCAGGATATGAT 1140
-----+-----+-----+-----+-----+-----+
CCATTAAGATCCTACCGGTTGTCGCTTCTGTACCCCTTGAAGATGAGGCGTCCTATACTA
G N S R M A N S E D M G N F Y S A G Y D -

1141 CCTCTCTTTTACGTCCACCATGCGAATGTGACAGGATGTGGCAAATCTGGAAAGATATT 1200
-----+-----+-----+-----+-----+-----+
GGAGAGAAAATGCAGGTGGTACGCTTACAGCTGTCTACACCGTTTAGACCTTTCTATAA
P L F Y V H H A N V D R M W Q I W K D I -

1201 GACAAGAAGACACACAAGGATCCGACCTCTGGCGACTGGCTAAATGCATCATACGTGTTT 1260
-----+-----+-----+-----+-----+-----+
CTGTTCTTCTGTGTGTTTCTAGGCTGGAGACCGCTGACCGATTTACGTAGTATGCACAAA
D K K T H K D P T S G D W L N A S Y V F -

1261 TACGATGAGAATGAAAATCTTGTACGTGTCTACAACCGAGACTGTGTAGACATTAATCGG 1320
-----+-----+-----+-----+-----+-----+
ATGCTACTCTTACTTTTAGAACATGCACAGATGTTGGCTCTGACACATCTGTAATTAGCC
Y D E N E N L V R V Y N R D C V D I N R -

1321 ATGGGATATGACTACGAAAGGTCAGCAATCCCATGGATCCGTAGTCGGCCGACTGCACAT 1380
-----+-----+-----+-----+-----+-----+
TACCCTATACTGATGCTTTCCAGTCGTTAGGGTACCTAGGCATCAGCCGCTGACGTGTA
M G Y D Y E R S A I P W I R S R P T A H -

1381 GCGAAGGGGGCGAACGTTGCTGCTAAGTCTGCTGGAATCGTGCAGAAGGTGGAGGATATC 1440
-----+-----+-----+-----+-----+-----+
CGCTTCCCCGCTTGCAACGACGATTACAGACGCTTAGCACGTCTCCACCTCCTATAG
A K G A N V A A K S A G I V Q K V E D I -

1441 GTATTCCCGCTGAAGTTAAACAAGATAGTGAAGGTTCTAGTGAAGAGGCCAGCTACAAAC 1500
-----+-----+-----+-----+-----+-----+
CATAAGGGCGACTTCAATTTGTTCTATCACTTCCAAGATCACTTCTCCGGTCGATGTTG
V F P L K L N K I V K V L V K R P A T N -

1501 AGGACCAAGGAGGGAAAGGAGAAAGCAAATGAGCTGTTGTTTCGTGAATGGAATCACGTTT 1560
-----+-----+-----+-----+-----+-----+
TCCTGGTTCCTCCCTTTCTCTTTCTGTTTACTCGACAACAAGCACTTACCTTAGTGCAA
R T K E G K E K A N E L L F V N G I T F -

1561 GATGCTGAGCGGTTTCTAAAGATTGACGTGTTTGTCAACGACGTCGACGATGGAATTCAG 1620
-----+-----+-----+-----+-----+-----+
CTACGACTCGCCAAAGATTTCTAAGTGCACAAACAGTTGCTGCAGCTGCTACCTTAAGTC
D A E R F L K I D V F V N D V D D G I Q -

1621 ACCACCGCTGCTGATAGTGAGTTTGTGTTAGTTTCGCACAGTTGCCACATAACCATGGC 1680
-----+-----+-----+-----+-----+-----+
TGGTGGCGACGACTATCACTCAAACGACCATCAAAGCGTGTCAACGTGTATTGTTACCCG
T T A A D S E F A G S F A Q L P H N H G -

FIG 1
(cont.)


```
1681 GACAAGATGTTTATGAGGAGTGGGGCAGCGTTCGGGATCACGGAGCTCTTGAAGACATT
-----+-----+-----+-----+-----+ 1740
CTGTTCTACAAATACTCCTCACCCCGTCGCAAGCCCTAGTGCCTCGAGAACCTTCTGTAA
D K M F M R S G A A F G I T E L L E D I -
1741 GAAGCTGAAGGTGATGACTCTGTTGTTGTGACATTGGTGCCGAGAACAGGGTGTGATGAA
-----+-----+-----+-----+-----+ 1800
CTTCGACTTCCACTACTGAGACAACAACACTGTAACCACGGCTCTGTCCCACACTACTT
E A E G D D S V V V T L V P R T G C D E -
1801 GTAAC TATTGGCGAGATCAAGATT CAGCTGGT TCCCAT TGT TTAAGTCTATTGAAGTAA
-----+-----+-----+-----+-----+ 1860
CATTGATAACCGCTCTAGTTCTAAGTCGACCAAGGGTAACAAATTCAGATAACTTCATT
V T I G E I K I Q L V P I V *
1861 TGCATTTTCAATTGTCATTAGTATGCATGGGTACGTAAATCTGTTTCGCTGTCTGGTTATC
-----+-----+-----+-----+-----+ 1920
ACGTAAAAGTTAACAGTAATCATACGTACCCATGCATTTAGACAAGCGACAGACCAATAG
1921 GAGGATTTTTGATGTTCTCGTAACCAAATAATAAGGATTGTCATTCCATGTTTGAATCG
-----+-----+-----+-----+-----+ 1980
CTCCTAAAACTACAAGAGCATTGGTTTATTATTCCCTAACAGTAAGGTACAAACCTTAGC
1981 TGTAACCGCAGGCATGCATATGTTTGATTGTTATTTTACTTGAAGCACTTCTGTTT TAG
-----+-----+-----+-----+-----+ 2040
ACATTGGCGTCCGTACGTATACAACTAACAATAAAAAATGAACTTCGTGAAGACAAAATC
2041 TAAAAAAAAAAAAAAAAA
-----+----- 2057
ATTTTTTTTTTTTTTTTT
```

FIG 1
(cont.)

b

1 CACGCCACCCTTCTCTCTCTCTCTCTCTGGTCTACTGAACAGTAATAGACATGTCCCT
-----+-----+-----+-----+-----+ 60
GTGCGGTGGGAAGAGAGAGAGAGAGAGAGACCAGATGACTTGTCATTATCTGTACAGGGA
T P P F S L S L S L V Y * T V I D M S L -

61 GCTGTTGAACTCTAGCTTCACCGGTGCTTCCTCTGCATGCCTCCTCCAACGGGAAAGGTC
-----+-----+-----+-----+-----+ 120
CGACAACTTGAGATCGAAGTGGCCACGAAGGAGACGTACGGAGGAGGTTGCCCTTTCCAG
L L N S S F T G A S S A C L L Q R E R S -

121 CCGCCGCCGCCCTCCACGTCCCTGGCGTGACATGCCGCCAGGGCAGTAATGGTGACCG
-----+-----+-----+-----+-----+ 180
GGCGGCCGCCGCCGAGGTGCAGGGACCGCACTGTACGGCGGTCCCGTCATTACCACTGGC
R R R R L H V P G V T C R Q G S N G D R -

181 CAGAGATGCCGCCCCCAGCAGCAGTCGCCGCCGCTGCTGGATCGGCGCGACATGCTGTT
-----+-----+-----+-----+-----+ 240
GTCTCTACGGCGGGGGGTCTGCTCGTCAGCGCGCGGACGACCTAGCCGCGCTGTACGACAA
R D A A P Q Q Q S P P L L D R R D M L L -

241 GGGTTTAGGAGGGCTTTACGGCGTGACCGCAGGACCCAAGGTTCTGGCGGCGCCGATAAT
-----+-----+-----+-----+-----+ 300
CCCAAATCCTCCCGAAATGCCGCACTGGCGTCTGGGTTCCAAGACCGCCGCGGTATTA
G L G G L Y G V T A G P K V L A A P I M -

301 GCCGCCGGATCTGTCCAAGTGCTACCCTGCCACCGCACCTGCCCTCGACAACAAATGCTG
-----+-----+-----+-----+-----+ 360
CGGCGGCCTAGACAGGTTACGATGGGACGGTGGCGTGACGGGAGCTGTTGTTTACGAC
P P D L S K C Y P A T A P A L D N K C C -

361 CCCGCCCTTACGACCCCGGCGAGACGATCTCGGAGTACAGCTTCCCTGCTACGCCCTCCG
-----+-----+-----+-----+-----+ 420
GGGCGGAATGCTGGGGCCGCTCTGCTAGAGCCTCATGTGAAGGGACGATGCGGGGAGGC
P P Y D P G E T I S E Y S F P A T P L R -

421 GGTGCGGCGGCCCGGCCCATATCGTGAAGGACGATCAGGAGTATATGGACAAGTACAAGGA
-----+-----+-----+-----+-----+ 480
CCACGCCGCCGCCGGGTATAGCACTTCTGCTAGTCCTCATATACCTGTTTCATGTTCT
V R R P A H I V K D D Q E Y M D K Y K E -

481 GGCAGTGAGGAGGATGAAGAATCTGCCGGCAGACCACCTTGGAACTACTACCAGCAGGC
-----+-----+-----+-----+-----+ 540
CCGTCACTCCTCCTACTTCTTAGACGGCCGTCTGGTGGGAACCTTGATGATGGTTCGTCG
A V R R M K N L P A D H P W N Y Y Q Q A -

541 GAACATCCACTGCCAGTATTGCAACTACGCCCTACCACCAGCAAAATACCGACGACGTGCC
-----+-----+-----+-----+-----+ 600
CTTGTAGGTGACGGTCATAACGTTGATGCGGATGGTGGTTCGTTTATGGCTGCTGCACGG
N I H C Q Y C N Y A Y H Q Q N T D D V P -

601 CATCCAGGTCCACTTCAGCTGGATCTTCCTCCCATGGCACCGCTACTACCTCCACTTCTA
-----+-----+-----+-----+-----+ 660
GTAGGTCCAGGTGAAGTCGACCTAGAAGGAGGGTACCGTGGCGATGATGGAGGTGAAGAT
I Q V H F S W I F L P W H R Y Y L H F Y -

661 CGAAAGGATCCTCGGCAAGCTCATCGACGACGACACCTTCACCATCCCATTCTGGAAGTC
-----+-----+-----+-----+-----+ 720
GCTTTCCTAGGAGCCGTTTCAGTAGCTGCTGTGGAAGTGGTAGGGTAAGACCTTGAC
E R I L G K L I D D D T F T I P F W N W -

721 GGACACCAAGGACGGGATGACGTTCCCCGCCATCTTCCAGGATGCGGCATCCCCGCTGTA
-----+-----+-----+-----+-----+ 780
CCTGTGGTTTCTGCCCTACTGCAAGGGCGGTAGAAGGTCTACGCCGTAGGGGCGACAT
D T K D G M T F P A I F Q D A A S P L Y -

781 CGACCCGAGACGCGACCAACGCCACGTCAAGGACGGCAAGATCCTCGACCTCAAGTACGC
-----+-----+-----+-----+-----+ 840
GCTGGGCTCTGCGCTGGTTGCGGTGCAAGTTCCTGCCGTTCTAGGAGCTGGAGTTCATGCG
D P R R D Q R H V K D G K I L D L K Y A -

FIG 2

	CTACACCAGAAACACTGCATCCGACAGCGAGATCATACGGGAGAACCTCTGCTTCATACA	
841	-----+-----+-----+-----+-----+	
	GATGTGGCTTTTGTGACGTAGGCTGTGCTCTAGTATGCCCTCTTGAGACGAAGTATGT	900
	Y T E N T A S D S E I I R E N L C F I Q -	
	GAAGACGTTCAAGCACAGCCTGTGCTGGCGGAAGTGTTCATGGGGGATCCCCTGCGCGC	
901	-----+-----+-----+-----+-----+	
	CTTCTGCAAGTTTCGTGTGCGACAGCGACCGCCTTGACAAGTACCCCCCTAGGGCACGCGG	960
	K T F K H S L S L A E L F M G D P V R A -	
	GGGGGAGAAGGAGATCCAGGAGGCTAATGGGCAGATGGAAGTCATCCACAATGCGGCGCA	
961	-----+-----+-----+-----+-----+	
	CCCCCTCTTCCTCTAGGTCCTCCGATTACCGCTACCTTCAGTAGGTGTACGCCGCGT	1020
	G E K E I Q E A N G Q M E V I H N A A H -	
	CATGTGGGTCGGAGAGCCGGACGGATACAAGGAAAACATGGGGGACTTCTCCACCGCCGC	
1021	-----+-----+-----+-----+-----+	
	GTACACCCAGCCTCTCGCCTGCCTATGTTCTTTGTACCCCTGAAGAGGTGGCGGCG	1080
	M W V G E P D G Y K E N M G D F S T A A -	
	CCGCGATTCTGTTTTCTTCTGCCACCATCCAATGTCGACCGCATGTGGGACATCTACCG	
1081	-----+-----+-----+-----+-----+	
	GGCGCTAAGACAAAAGAAGACGGTGGTAAGGTTACAGCTGGCGTACACCCTGTAGATGGC	1140
	R D S V F F C H H S N V D R M W D I Y R -	
	CAACCTCCGCGCAACCGCGTCGAGTTCAAGACAACGACTGGTTGGACAGCACCTTCCT	
1141	-----+-----+-----+-----+-----+	
	GTTGGAGGCGCGTGGCGCAGCTCAAGCTTCTGTTGCTGACCAACCTGTCGTGGAAGGA	1200
	N L R G N R V E F E D N D W L D S T F L -	
	CTTCCACGACGAGAACGAACAGCTCGTCAAAGTCAAGATGAGCGACTGCCTCAACCCGAC	
1201	-----+-----+-----+-----+-----+	
	GAAGGTGCTGCTCTTGCTTGTGCGAGCAGTTTCACTTCTACTCGCTGACGGAGTTGGGCTG	1260
	F H D E N E Q L V K V K M S D C L N P T -	
	CAAGCTTCGGTACACGTTTCGAGCAAGTTCCCTCCCATGGCTGGGCAAAATCAATTGCCA	
1261	-----+-----+-----+-----+-----+	
	GTTCAAGCCATGTGCAAGCTCGTTCAAGGGGAGGGTACCGACCCGTTTTAGTTAACGGT	1320
	K L R Y T F E Q V P L P W L G K I N C Q -	
	GAAGACGGCAGAGACGAAGTCCAAGGCCACGACGGAGCTGTGCTGACGCGCGTGAACGA	
1321	-----+-----+-----+-----+-----+	
	CTTCTGCCGCTCTGCTTCAGGTTCCGGTGCTGCCTCGACAGCGACTGCGCGCACTTGCT	1380
	K T A E T K S K A T T E L S L T R V N E -	
	ATTCGGGACGACGGCCCAGGCACTCGACGCGAGCAACCCGCTGCGGGTGATCGTGGAAG	
1381	-----+-----+-----+-----+-----+	
	TAAGCCCTGCTGCCGGGTCCGTGAGCTGCGCTCGTTGGGCGACGCCCACTAGCACCGTTC	1440
	F G T T A Q A L D A S N P L R V I V A R -	
	GCCGAAGAAGAACCGBAAGAAGAAGGAGAAGCAAGAGAAGGTGGGGGTGATTCAGATCAA	
1441	-----+-----+-----+-----+-----+	
	CGGCTTCTTCTTGGCGTTCTTCTTCTTCTTCTTCCACCCCACTAAGTCTAGTT	1500
	P K K N R K K K E K Q E K V G V I Q I K -	
	GGATATTAAGGTGACCACCAACGAGACAGCTCGCTTCGACGCTCTATGTCGCGGTTCTTTA	
1501	-----+-----+-----+-----+-----+	
	CCTATAATTCCACTGGTGGTTGCTCTGTGAGCGAAGCTGCAGATACAGCGCAAGGAAT	1560
	D I K V T T N E T A R F D V Y V A V P Y -	
	CGGTGACCTCGCCGGACCCGACTACGGCGAGTTCGCGGGCAGCTACGTGAGGCTGGCGCA	
1561	-----+-----+-----+-----+-----+	
	GCCACTGGAGCGGCTGGGCTGATGCCGCTCAAGCGCCCGTCGATGCACTCCGACCGCGT	1620
2	G D L A G P D Y G E F A G S Y V R L A H -	
t.)	TAGGATGAAGGGAAGCGACGGGACCGAAAAAGCAGGGCCCCAAGAAGAAGGAAAACTCAA	
1621	-----+-----+-----+-----+-----+	
	ATCCTACTTCCCTTCGCTGCCCTGGCTTTTTCGTCGGGGGTTCTTCTTCCCTTTTGAGTT	1680
	R M K G S D G T E K Q G P K K K G K L K -	

1681 GCTGGGTATTACGCCGCTGCTCGAGGACATCGATGCTGAGGACGCCGACAAGTTGGTGGT
-----+-----+-----+-----+ 1740
CGACCCATAATGCGGCGACGAGCTCCTGTAGCTACGACTCCTGCGGCTGTTCAACCACCA
L G I T P L L E D I D A E D A D K L V V -
1741 CACCCCTGGTTCTCCGCACTGGGAGCGTCACCGTGGGGGGAGTTTCCATCAATCTCCTGCA
-----+-----+-----+-----+ 1800
GTGGGACCAAGAGGCGTGACCCTCGCAGTGGCACCCCTCAAAGGTAGTTAGAGGACGT
T L V L R T G S V T V G G V S I N L L Q -
1801 GACAGATTCTACCGCCGCCATCTAAATGATGGCCTCGGATCACAGCTTCTCCCCGCTTAA
-----+-----+-----+-----+ 1860
CTGTCTAAGATGGCGGCGGTAGATTTACTACCGGAGCCTAGTGTGCGAAGAGGGGCGAATT
T D S T A A I *
1861 GTTGGAGTGATCGATTACTGGTGCTGCTTTCTTCCCTGTCGTTCTTGCTATCTTCTT
-----+-----+-----+-----+ 1920
CAACCTCACTAGCTAATGACCACGACGAAAGAAGGAGGGACAGCAAGAACGATAGAAGAA
1921 GATCTGGAACGATCCTTCAATAATTAGGGCATGACAGTAGTCGTCGCCCCGATCCCATATG
-----+-----+-----+-----+ 1980
CTAGACCTTGCTAGGAAGTTATTAATCCCGTACTGTCATCAGCAGCGGGCTAGGGTATAC
1981 TACGTGTTGGTCTCAACAGCTGTACATGTGACGTTATGGTGTGACTATATATTTTATTGC
-----+-----+-----+-----+ 2040
ATGCACAACCAGAGTTGTCGACATGTACTGCAATACCACACTGATATATAAAATAACG
2041 GGTCATCCTTGTTTCTTTCTTAAAAA
-----+-----+-----+ 2078
CCAGTAGGAACAAAGAAAGAAATTTTTTTTTTTTTTTTTTT

FIG 2
(cont.)

a

1 AATGTGGATCGGATGTGGACGGTGTGGAAGAAGCTGCACGGCGACAAGCCGGAGTTCGTC
-----+-----+-----+-----+-----+ 60
TTACACCTAGCCTACACCTGCCACACCTTCTTCGACGTGCCGCTGTTCGGCCTCAAGCAG
N V D R M W T V W K K L H G D K P E F V -

61 GACCAGGAGTGGCTCGAGTCTGAATTCACCTTCTACGACGAGAATGTGCGCCTGCGCAGG
-----+-----+-----+-----+-----+ 120
CTGGTCCTCACCGAGCTCAGACTTAAGTGAAGATGCTGCTCTTACACGCGGACGCGTCC
D Q E W L E S E F T F Y D E N V R L R R -

121 ATCAAGGTGCGCGACGTGTGAACATAGACAAACTCAGGTACCGGTACGAAGACATCGAC
-----+-----+-----+-----+-----+ 180
TAGTTCCACGCGCTGCACAACCTTGTATCTGTTTGAGTCCATGGCCATGCTTCTGTAGCTG
I K V R D V L N I D K L R Y R Y E D I D -

181 ATGCCATGGCTCGCTGCACGTCCCAAGCCTTCCGTTACCCCTAAGATCGCGCGCGACATA
-----+-----+-----+-----+-----+ 240
TACGGTACCGAGCGACGTGCAGGGTTCCGAAGGCAAGTGGGATTCTAGCGCGCGCTGTAT
M P W L A A R P K P S V H P K I A R D I -

241 TTGAAGAAGCGTAATGGCGAAGGCGTACTGAGAATGCCCGGCGAAACGGATCGTTCACAA
-----+-----+-----+-----+-----+ 300
AACTTCTTCGCATTACCGCTTCCGCATGACTCTTACGGGCCGCTTTGCCTAGCAAGTGTT
L K K R N G E G V L R M P G E T D R S Q -

301 CTCTCCGAAGATGGTAGCTGGACACTGGACAAGAGCATCACCGTGAGGGTTGACAGGCCA
-----+-----+-----+-----+-----+ 360
GAGAGGCTTCTACCATCGACCTGTGACCTGTTCTCTGTAGTGGCACTCCCAACTGTCCGGT
L S E D G S W T L D K S I T V R V D R P -

361 AGGATCAACAGGACAGGGCAAGAAAAGAGGAAGAAGAGGAGATCTTATTGGTCTACGGA
-----+-----+-----+-----+-----+ 420
TCCTAGTTGTCTGTCCCGTTCTTTTCTCCTTCTTCTCTCTAGATAACCAGATGCCT
R I N R T G Q E K E E E E E I L L V Y G -

421 ATCGATACTAAGAGAAGCAGATTCTGTCAAATTCGATGTGTTTCATCAACGTCGTCGACGAA
-----+-----+-----+-----+-----+ 480
TAGCTATGATTCTCTTCGTCTAAGCAGTTTAAGCTACACAAGTAGTTGCAGCAGCTGCTT
I D T K R S R F V K F D V F I N V V D E -

481 ACCGTGCTGAACCCAAAGTCGAGGGAGTTTCGACGGGACCTTCGTCAATCTCCACCACGTC
-----+-----+-----+-----+-----+ 540
TGGCAGCACTTGGGTTTTAGCTCCCTCAAGCGTCCCTGGAAGCAGTTAGAGGTGGTGCAG
T V L N P K S R E F A G T F V N L H H V -

541 TCGAGGACGAAAAGCCATGAGGATGGCGGCGTGGGTTTCAAGATGAAAAGCCACCTTAAG
-----+-----+-----+-----+-----+ 600
AGCTCCTGCTTTTGGTACTCCTACCGCCGACCCAAGCTTCTACTTTTCGGTGGGAATTC
S R T K S H E D G G V G S K M K S H L K -

601 CTCGGTATATCGGAACCTTTTGAAGACCTCGAGGCAGACGAAGATGATTGCATCTGGGTG
-----+-----+-----+-----+-----+ 660
GAGCCATATAGCCTTGAAAACCTTCTGGAGCTCCGTCTGCTTCTACTAACGTAGACCCAC
L G I S E L L E D L E A D E D D C I W V -

661 ACACTGGTGCCAAGAGGCGGCACGGGGGTCAACACCACCGTAGACGGCGTCCGGATCGAC
-----+-----+-----+-----+-----+ 720
TGTGACCACGGTTCTCCGCCGTGCCCCCAGTTGTGGTGGCATCTGCCGAGGCCTAGCTG
T L V P R G G T G V N T T V D G V R I D -

721 TACATGAAGTAGTGAACCGGCACGCCCTCCTCCCCCTCCCCATCAGAAGTGGTATAATAT
-----+-----+-----+-----+-----+ 780
ATGTACTTCATCACTTGGCCGTGCGGCGAGGAGGGGAGGGGTAGTCTTACCATATTATA
Y M K *

FIG 3

781 TTATATTGGATCGAGGCTCGTGGTATCTTTTGATAAGAGTAAGTTCCATAAATTTAGAAG
-----+-----+-----+-----+-----+-----+ 840
AATATAACCTAGCTCCGAGCACCATAGAAACTATTCTCATTCAAGGTATTTAAATCTTC
841 AAGAATCATGTTCTTTATTTATATTAAATCAATGTGATTTGTCCAAAAAAAAAAAAAAAA
-----+-----+-----+-----+-----+-----+ 900
TTCTTAGTACAAGAAATAAATATAATTTAGTTACACTAAACAGGTTTTTTTTTTTTTTTTT

FIG 3
(cont.)

A. CLASSIFICATION OF SUBJECT MATTERInt Cl⁶: C12N 15/53, 15/29, 5/04; A01H 5/00, 1/00

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC6: C12N, A01H. Chemical Abstracts. All through Electronic Databases

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched
Biotechnology Abstracts Through Electronic Databases

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

DERWENT DATABASES: WPAT & JAPIO Search terms: polyphenol(oxidase#, poly(phenol(oxidase#, ppo#, catechol(oxidase#, tryosinase#, diphenol(oxidase#, phenolase#, diphenolase#, mono(phenol(mono(oxygenase#, monophenol(mono(oxygenase#, brown:, C12N-015/IC, A01H/IC.

"BIOT" Search terms: EC-1.10.3.1 and A1/CL

"CASM" Search terms: The primers of claims 4, 5, 9, 10 and 11 were searched as partial nucleotide sequences. Also searched were polyphenol(oxidase#, diphenol(oxidase#, phenolase#, diphenolase#, gene#.

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X Y	WO 93/2195 (COMMONWEALTH SCIENTIFIC AND INDUSTRIAL RESEARCH ORGANISATION) 4 February 1993 Int Cl ⁵ : C12N 15/53, 9/02 See entire document	1-7 2,22,23
X	I.B.Dry & S.P. Robinson: "Molecular cloning and characterisation of grape berry polyphenol oxidase". Plant Molecular Biology, Vol 26, pp 495-502, 1994 See entire document	1,2

☒ Further documents are listed in the continuation of Box C☒ See patent family annex

* Special categories of cited documents:

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"X"

document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

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document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

"&"

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Date of the actual completion of the international search

18 June 1996

Date of mailing of the international search report

02 JUL 1996

Name and mailing address of the ISA/AU
AUSTRALIAN INDUSTRIAL PROPERTY ORGANISATION
PO BOX 200
WODEN ACT 2606
AUSTRALIA Facsimile No.: (06) 285 3929

Authorized officer

ROBYN PORTER

Telephone No.: (06) 283 2318

C (Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	Derwent BIOT Online abstract Accession No. 90-12612. Abstracts of the Annual Meeting of the American Society for Microbiology, 90 th Meeting, page 163, 1990. Williams et al: "Molecular cloning and partial characterisation of the polyphenol-oxidase (PPO) gene of <i>Coriolus versicolor</i> ". See entire abstract	1
X	WO 88/02372 (DONALD GUTHRIE FOUNDATION FOR MEDICAL RESEARCH), 7 April 1988, Int Cl ⁴ : C07H 15/12, C12Q 1/68, C12P 21/02, C12Q 1/26, C12P 13/22, C12N 15/00, 7/00.	1
Y	See pages 1, 20-21, claims 1, 6, 7.	2
Y	Derwent BIOT Online Abstract Accession No. 95-05853. Abstracts of Papers of the American Chemical Society, 208 th Meeting, Part 1, AGFD107, 1994. Steffans: "Modification of polyphenol-oxidase expression in crop plants". See entire abstract.	20,21
Y	Derwent BIOT Online Abstract Accession No. 95-05846. Abstracts of Papers of the American Chemical Society, 208 th Meeting, Part 1, AGFD2, 1994. Martinez and Whitaker: "Isolation of the gene encoding grape polyphenol-oxidase and study of PPO control with antisense RNA". See entire abstract	20,21
Y	WO 94/03607 (KEYGENE N.V.) 17 February 1994, Int Cl ⁵ : C12N 15/53, 15/11, A01H 5/00. See page 1 lines 3-11, claims 1, 15, 16.	22,23
Y	WO 93/15599 (CORNELL RESEARCH FOUNDATION), 19 August 1993, Int Cl ⁵ : A01H 5/00, C12N 15/00, 15/29. See claims 2, 3	22,23

This Annex lists the known "A" publication level patent family members relating to the patent documents cited in the above-mentioned international search report. The Australian Patent Office is in no way liable for these particulars which are merely given for the purpose of information.

Patent Document Cited in Search Report				Patent Family Member			
WO	9302195	AU	23316/92	CA	2112998	EP	599868
		JP	7501686	NZ	243594		
WO	8802372	AU	81547/87	CA	1293940	EP	290504
		US	4898814				
WO	9403607	EP	606454	JP	7503376		
WO	9215599	EP	640613	JP	4275295	US	5470962
END OF ANNEX							